WHAT IS CLAIMED IS:

1	1. A method of detecting cancer cells in a biological sample from a			
2	mammal, the method comprising steps of:			
3	(i) providing the biological sample from the mammal; and			
4	(ii) detecting a nucleic acid molecule encoding a PRC17 polypeptide			
5	comprising at least 85% amino acid sequence identity to an amino acid sequence of SEQ			
6	ID NO:2 or at least 70% amino acid identity to an amino acid sequence of SEQ ID NO:4			
7	or SEQ ID NO:6 in the biological sample, wherein an increase in the level of the nucleic			
8	acid molecule in the sample compared to normal indicates the presence of cancer cells.			
1	2. The method of claim 1, wherein the polypeptide has an amino acid			
2 -	sequence of SEQ ID NO:2, SEQ ID NO:4 or SEQ ID NO:6.			
1	3. The method of claim 1, wherein the detecting step further			
2	comprises:			
3	(a) contacting the nucleic acid molecule with a probe under conditions in			
4	which the probe selectively hybridizes to the nucleic acid molecule to form a stable			
5	hybridization complex; and			
6	(b) detecting the hybridization complex.			
1	4. The method of claim 3, wherein the contacting step further			
2	comprises a step of amplifying the gene in an amplification reaction.			
1	5. The method of claim 4, wherein the amplification reaction is a			
2	polymerase chain reaction.			
1	6. The method of claim 1, wherein the nucleic acid is an mRNA.			
1	7. The method of claim 1, wherein the biological sample is a tissue			
2	biopsy.			
1	8. The method of claim 7, wherein the cancer cells are selected from			
2	the group consisting of prostate tissue, breast tissue, lung tissue, and ovarian tissue.			
1	9. The method of claim 1, wherein the mammal is a human.			

1	10. A method of detecting a presence of cancer cells in a biological		
2	sample from a mammal, the method comprising steps of:		
3	(i) providing the biological sample from the mammal; and		
4	(ii) detecting an overexpression of a polypeptide comprising polypeptide		
5	comprising at least 85% amino acid sequence identity to an amino acid sequence of SEQ		
6	ID NO:2 or at least 70% amino acid identity to an amino acid sequence of SEQ ID NO:4		
7	or SEQ ID NO:6 in the biological sample, thereby detecting the presence of cancer cells		
8	in the biological sample.		
1	11. The method of claim 10, wherein the polypeptide has an amino		
2	acid sequence of SEQ ID NO:2, SEQ ID NO:4 or SEQ ID NO:6.		
L	12. The method of claim 10, wherein the polypeptide is detected using ar		
2	antibody that selectively binds to the polypeptide.		
l	13. The method of claim 10, wherein the biological sample is a tissue		
2	biopsy.		
1	14. The method of claim 10, wherein the cancer cells are selected from		
2	the group consisting of prostate cancer cells, breast cancer cells, lung cancer cells, and		
3	ovarian cancer cells.		
1	15. The method of claim 10, wherein the mammal is a human.		
1	16. A method of monitoring the efficacy of a therapeutic treatment of a		
2	cancer, the method comprising the steps of:		
3	(i) providing a biological sample from a mammal undergoing the		
4	therapeutic treatment; and		
5	(ii) detecting a level of a polypeptide comprising at least 85% amino acid		
6	sequence identity to an amino acid sequence of SEQ ID NO:2 or at least 70% amino acid		
7	identity to an amino acid sequence of SEQ ID NO:4 or SEQ ID NO:6 in the biological		
8	sample compared to a level in a biological sample from the mammal prior to, or earlier in		
9	the therapeutic treatment, thereby monitoring the efficacy of the therapy.		
1	17. The method of claim 16, wherein the polypeptide has an amino		
2	acid sequence of SEQ ID NO:2, SEQ ID NO:4 or SEQ ID NO:6.		

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1	18. The method of claim 16, wherein the cancer is selected from the		
2	group consisting of prostate cancer, ovarian cancer, lung cancer, and breast cancer.		
1	19. The method of claim 16, wherein the polypeptide is detected using		
2	an antibody that selectively binds to the polypeptide.		
1	20. A method of monitoring the efficacy of a therapeutic treatment of a		
2	· · ·		
3	cancer, the method comprising the steps of:		
	(i) providing a biological sample from a mammal undergoing the		
4	therapeutic treatment; and		
5	(ii) detecting a nucleic acid molecule encoding a PRC17 polypeptide		
6	comprising at least 85% amino acid sequence identity to an amino acid sequence of SEQ		
7	ID NO:2 or at least 70% amino acid identity to an amino acid sequence of SEQ ID NO:4		
8	or SEQ ID NO:6 in the biological sample compared to a level in a biological sample from		
9	the mammal prior to, or earlier in, the therapeutic treatment, thereby monitoring the		
10	efficacy of the therapy.		
1	21. An isolated nucleic acid encoding a PRC17 polypeptide, the		
2	nucleic acid encoding a polypeptide comprising at least 85% amino acid identity to an		
3	amino acid sequence of SEQ ID NO:2 or at least 70% identity to an amino acid sequence		
	•		
4	of SEQ ID NO:4 or SEQ ID NO:6.		
1	22. The nucleic acid of claim 21, wherein the nucleic acid encodes a		
2	PRC17 polypeptide that specifically binds to polyclonal antibodies generated against an		
3	amino acid sequence of SEQ ID NO:2, SEQ ID NO:4 or SEQ ID NO:6.		
1	23. The nucleic acid of claim 21, wherein the nucleic acid encodes a		
2	PRC17 polypeptide comprising an amino acid sequence of SEQ ID NO:2, SEQ ID NO:4		
3	or SEQ ID NO:6.		
1	24. The nucleic acid of claim 23, wherein the nucleic acid comprises a		
2	nucleotide sequence of SEQ ID NO:1, SEQ ID NO:3 or SEQ ID NO:5.		
1	25. The nucleic acid of claim 21, wherein the nucleic acid is amplified		

by primers that specifically hybridize under stringent hybridization conditions to a nucleic

acid having a nucleotide sequence of SEQ ID NO:1, SEQ ID NO:3 or SEQ ID NO:5.

cell or cell membrane.

1		26.	The nucleic acid of claim 21, wherein the nucleic acid specifically		
2	hybridizes under stringent hybridization conditions to a nucleic acid having a nucleotide				
3	sequence of S	SEQ II	NO:1, SEQ ID NO:3 or SEQ ID NO:5.		
1		27.	An isolated PRC17 polypeptide, the polypeptide comprising at		
2	least 85% am	ino aci	d sequence identity to an amino acid sequence of SEQ ID NO:2 or at		
3	least 70% amino acid identity to an amino acid sequence of SEQ ID NO:4 or SEQ ID				
4	NO:6.				
1		28.	The isolated polypeptide of claim 8, wherein the polypeptide		
2	specifically binds to polyclonal antibodies generated against SEQ ID NO:2, SEQ ID				
3	NO:4 or SEQ ID NO:6.				
1		29.	The isolated polypeptide of claim 8, wherein the polypeptide has		
2	an amino acid sequence of SEQ ID NO:2, SEQ ID NO:4 or SEQ ID NO:6.				
1		30.	An antibody that selectively binds to the polypeptide of claim 8.		
1		31.	An expression vector comprising the nucleic acid of claim 1.		
1		32.	A host cell transfected with the vector of claim 31.		
1		33.	A method of identifying a compound that modulates activity of a		
2	PRC17 polyp	eptide,	, the method comprising steps of:		
3		(i) co	ontacting the polypeptide with the compound, wherein the polypeptide		
4	comprises at	least 8:	5% amino acid sequence identity to an amino acid sequence of SEQ		
5	ID NO:2 or at least 70% amino acid identity to an amino acid sequence of SEQ ID NO:4				
6	or SEQ ID NO:6; and				
7		(ii) d	etermining the functional effect of the compound on the polypeptide.		
1	,	34.	The method of claim 33, wherein the polypeptide is linked to a		
2	solid phase.				
1		35.	The method of claim 33, wherein the polypeptide is expressed in a		

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1	36.	The method of claim 33, wherein the polypeptide has an amino		
2	acid sequence of SEQ	ID NO:2, SEQ ID NO:4 or SEQ ID NO:6.		
1	37.	A method of treating a disease or condition associated with the		
2	activity of a PRC17 p	olypeptide, the method comprising the step of administering to a		
3	subject a therapeutically effective amount of a compound identified using the method of			
4	claim 33.			
1	38.	The method of claim 37, wherein the subject is a human.		
1	39.	The method of claim 18, wherein the compound is an antibody.		
1	40.	A method of inhibiting proliferation of a cancer cell that expresses a		
2	polypeptide comprisin	g at least 85% amino acid sequence identity to an amino acid sequence		
3	of SEQ ID NO:2 or at	least 70% amino acid identity to an amino acid sequence of SEQ ID		
4	NO:4 or SEQ ID NO:6	6, the method comprising the step of contacting the cancer cell with a		
5	therapeutically effective	ve amount of an inhibitor of the polypeptide.		
1	41.	The method of claim 40, wherein the polypeptide has an amino acid		
2	sequence of SEQ ID N	O:2, SEQ ID NO:4 or SEQ ID NO:6.		
1	42.	The method of claim 40, wherein the cancer cell is selected from		
2	the group consisting of a prostate cancer cell, a breast cancer cancer cell, a lung cancer			
3	cell or an ovarian cancer cell.			